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SEQUENCE LISTING

<110> Cambridge Antibody Technology Limited Thompson, Julia E Lennard, Simon N Wilton, Alison J Braddock, Peta SH Du Fou, Sarah L McCafferty, John G Conroy, Louise A Tempest, Philip R <120> Specific binding members for TGF beta 1 <130> 28111/37903 <140> To be assigned <141> Herewith <150> PCT/GB00/01679 <151> 2000-05-02 <150> US 60/131,983 <151> 1999-04-30 <160> 25 <170> PatentIn Ver. 2.1 <210> 1 <211> 369 <212> DNA <213> Homo sapiens caggtgcagc tggtgcagtc tgggggaggc gtggtccagc ctgggaggtc cctgagactc 60 tectgtgcag cetetggatt cacettcagt agetatggca tgcactgggt eegecagget 120 ccaggcaagg ggctggagtg ggtggcagtt atatcatatg atggaagtat taaatactat 180 gcagactccg tgaagggccg attcaccatc tccagagaca attccaagaa cacgctgtat 240 ctgcaaatga acagcctgag agctgaggac acggctgtgt attactgtgc gcgaactggt 300 gaatatagtg gctacgatac gagtggtgtg gagctctggg ggcaagggac cacggtcacc 360 gtctcctca 369 <210> 2 <211> 123 <212> PRT <213> Homo sapiens <400> 2 Gln Val Gln Leu Val Gln Ser Gly Gly Val Val Gln Pro Gly Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr

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Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val 35 40 45

Ala Val Ile Ser Tyr Asp Gly Ser Ile Lys Tyr Tyr Ala Asp Ser Val
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Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr 65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
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Ala Arg Thr Gly Glu Tyr Ser Gly Tyr Asp Thr Ser Gly Val Glu Leu 100 105 110

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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr 20 25 30

Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Glu Leu Glu Trp Val 35 40 45

Ala Val Ile Ser Tyr Asp Gly Ser Ile Lys Tyr Tyr Ala Asp Ser Val
50 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr 65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys

85 90 95

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Leu Gly Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Ile Leu Leu Ile 35 40 45

Tyr Gly Thr Ser Thr Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly 50 60

Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Asn Ser Leu Gln Pro 65 70 75 80

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Thr Phe Gly Gly Gly Thr Arg Leu Glu Ile Lys

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gggaaagccc ctatcctcct gatctatggt acatccactt tacaaagtgg ggtcccgtca 180
aggttcagcg gcagtggatc tggcacagat ttcactctca ccatcaacag cctgcagcct 240
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Asp Arg Val Thr Ile Thr Cys Arg Ser Ser Gln Gly Ile Gly Asp Asp 20 25 30

Leu Gly Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Ile Leu Leu Ile 35 40 45

Tyr Gly Thr Ser Thr Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly 50 55 60

Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Asn Ser Leu Gln Pro 65 70 75 80

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ccaggcaagg agctggagtg ggtggcagtt atatcatatg atggaagtat taaatactat 180
gcagactccg tgaagggccg attcaccatc tccagagaca attccaagaa cacgctgtat 240

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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
20 25 30

Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Glu Leu Glu Trp Val 35 40 45

Ala Val Ile Ser Tyr Asp Gly Ser Ile Lys Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys 85 90 95

Ala Arg Thr Gly Glu Tyr Ser Gly Tyr Asp Thr Pro Ala Ser Pro Asp 100 105 110

Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser

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Gly

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Leu Gln Asp Ser Asn Tyr Pro Leu Thr
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<212> DNA
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<212> DNA
<213> Artificial Sequence
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<212> DNA
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